

1600

RAW SEQUENCE LISTING DATE: 03/19/2003
PATENT APPLICATION: US/09/446,634A TIME: 12:59:24

Input Set: A:\09-446,634 Sequence Listing.txt
Output Set: N:\CRF4\03192003\I446634A.raw

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3 <110> APPLICANT: Ono Pharmaceutical Co., Ltd.
      5 <120> TITLE OF INVENTION: NOVEL PLASMID DNA COMPRISING REPORTER GENE DNA AND USE OF
THE
              SAME
    . 8 <130> FILE REFERENCE: Q57282
     10 <140> CURRENT APPLICATION NUMBER: 09/446,634A
     11 <141> CURRENT FILING DATE: 1999-12-23
     13 <150> PRIOR APPLICATION NUMBER: JP 9-171440
     14 <151> PRIOR FILING DATE: 1997-06-27
                                                                 ENTERED
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     18 <170> SOFTWARE: PatentIn version 3.2
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    26 <223> OTHER INFORMATION: Primer F1
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	<220> FEATURE:		
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146	<223> OTHER INFORMATION: Primer F6	
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Input Set : A:\09-446,634 Sequence Listing.txt
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217 <211> LENGTH: 38 218 <212> TYPE: DNA 219 <213> ORGANISM: Artificial Sequence 221 <220> FEATURE: 222 <223> OTHER INFORMATION: Multiple cloning site DNA linker - sense strand 6 224 <400> SEQUENCE: 17 225 gaattcgtcg acggtaccga tatcgagctc gcggccgc 38 228 <210> SEQ ID NO: 18 229 <211> LENGTH: 85 230 <212> TYPE: DNA 231 <213> ORGANISM: Artificial Sequence 233 <220> FEATURE: 234 <223> OTHER INFORMATION: Gal4 responsive element 236 <400> SEQUENCE: 18 237 tegaeggagt actgteetee gegaeggagt aetgteetee gegaeggagt aetgteetee 60 239 gcgacggagt actgtcctcc gagct 85 242 <210> SEQ ID NO: 19 243 <211> LENGTH: 20 244 <212> TYPE: DNA 245 <213> ORGANISM: Artificial Sequence 247 <220> FEATURE: 248 <223> OTHER INFORMATION: Gal4 responsive element 250 <400> SEQUENCE: 19 20 251 cgacggagta ctgtcctccg 254 <210> SEQ ID NO: 20 255 <211> LENGTH: 9 256 <212> TYPE: PRT 257 <213> ORGANISM: Artificial Sequence 259 <220> FEATURE: 260 <223> OTHER INFORMATION: Influenza hemagglutinin epitope 262 <400> SEQUENCE: 20 264 Tyr Pro Tyr Asp Val Pro Asp Tyr Ala 268 <210> SEQ ID NO: 21 269 <211> LENGTH: 38 270 <212> TYPE: DNA 271 <213> ORGANISM: Artificial Sequence 273 <220> FEATURE: 274 <223> OTHER INFORMATION: Multiple cloning site DNA linker - antisense strand 276 <400> SEQUENCE: 21 277 gcggccgcga gctcgatatc ggtaccgtcg acgaattc 38 280 <210> SEQ ID NO: 22 281 <211> LENGTH: 335 282 <212> TYPE: PRT 283 <213> ORGANISM: Homo sapiens 286 <220> FEATURE: 287 <221> NAME/KEY: mat peptide 288 <222> LOCATION: (17)..(355)

290 <400> SEQUENCE: 22

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292 Met Leu Gly Ile Trp Thr Leu Leu Pro Leu Val Leu Thr Ser Val Ala
                           -10
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296 Arg Leu Ser Ser Lys Ser Val Asn Ala Gln Val Thr Asp Ile Asn Ser
                                        10
300 Lys Gly Leu Glu Leu Arg Lys Thr Val Thr Thr Val Glu Thr Gln Asn
                                    25
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304 Leu Glu Gly Leu His His Asp Gly Gln Phe Cys His Lys Pro Cys Pro
308 Pro Gly Glu Arg Lys Ala Arg Asp Cys Thr Val Asn Gly Asp Glu Pro
312 Asp Cys Val Pro Cys Gln Glu Gly Lys Glu Tyr Thr Asp Lys Ala His
                       70
                                            75
316 Phe Ser Ser Lys Cys Arg Arg Cys Arg Leu Cys Asp Glu Gly His Gly
320 Leu Glu Val Glu Ile Asn Cys Thr Arg Thr Gln Asn Thr Lys Cys Arg
                                   105
               100
324 Cys Lys Pro Asn Phe Phe Cys Asn Ser Thr Val Cys Glu His Cys Asp
          115
                               120
328 Pro Cys Thr Lys Cys Glu His Gly Ile Ile Lys Glu Cys Thr Leu Thr
329 130
                            135
                                                140
332 Ser Asn Thr Lys Cys Lys Glu Glu Gly Ser Arg Ser Asn Leu Gly Trp
                       150
                                            155
336 Leu Cys Leu Leu Leu Pro Ile Pro Leu Ile Val Trp Val Lys Arg
                                        170
                   165
340 Lys Glu Val Gln Lys Thr Cys Arg Lys His Arg Lys Glu Asn Gln Gly
               180
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344 Ser His Glu Ser Pro Thr Leu Asn Pro Glu Thr Val Ala Ile Asn Leu
           195
                                200
                                                    205
348 Ser Asp Val Asp Leu Ser Lys Tyr Ile Thr Thr Ile Ala Gly Val Met
                           215
                                                220
352 Thr Leu Ser Gln Val Lys Gly Phe Val Arg Lys Asn Gly Val Asn Glu
                       230
                                            235
356 Ala Lys Ile Asp Glu Ile Lys Asn Asp Asn Val Gln Asp Thr Ala Glu
                   245
                                        250
360 Gln Lys Val Gln Leu Leu Arg Asn Trp His Gln Leu His Gly Lys Lys
               260
                                    265
364 Glu Ala Tyr Asp Thr Leu Ile Lys Asp Leu Lys Lys Ala Asn Leu Cys
           275
                                280
368 Thr Leu Ala Glu Lys Ile Gln Thr Ile Ile Leu Lys Asp Ile Thr Ser
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382 <220> FEATURE:
383 <221> NAME/KEY: mat_peptide
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VERIFICATION SUMMARY

DATE: 03/19/2003

PATENT APPLICATION: US/09/446,634A

TIME: 12:59:25

Input Set : A:\09-446,634 Sequence Listing.txt
Output Set: N:\CRF4\03192003\I446634A.raw